RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

09/936,271 -C
IFW16
5/9/5

ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 05/09/2005
PATENT APPLICATION: US/09/936,271C TIME: 17:04:59

Input Set : A:\MTS3USA.seq.txt

3 <110> APPLICANT: Yousef, George M.

Output Set: N:\CRF4\05092005\I936271C.raw

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Diamandis, Eleftherios
 6 <120> TITLE OF INVENTION: Novel Human Kallikrein-Like Genes
 8 <130> FILE REFERENCE: MTS3USA
10 <140> CURRENT APPLICATION NUMBER: US 09/936,271C
11 <141> CURRENT FILING DATE: 2001-09-10
13 <150> PRIOR APPLICATION NUMBER: PCT/CA00/00258
14 <151> PRIOR FILING DATE: 2000-03-09
16 <150> PRIOR APPLICATION NUMBER: US 60/124,260
17 <151> PRIOR FILING DATE: 1999-03-11
19 <150> PRIOR APPLICATION NUMBER: US 60/127,386
20 <151> PRIOR FILING DATE: 1999-04-01
22 <150> PRIOR APPLICATION NUMBER: US 60/144,919
23 <151> PRIOR FILING DATE: 1999-07-21
25 <160> NUMBER OF SEQ ID NOS: 97
27 <170> SOFTWARE: PatentIn version 3.2
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 4740
31 <212> TYPE: DNA
32 <213> ORGANISM: Homo sapiens
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                                                                         120
39 getgecagee cettetggge ceccaaceae tgeetggtea gagttgagge ageetgagag
                                                                         180
41 agttgagetg gaagtttgea geacetgace eetggaacae ateceetggg ggeaggeeag
                                                                         240
43 cccaggctga ggatgcttat aagccccaag gaggcccctg cggaggcagc aggctggagc
                                                                         300
45 tcagcccagc agtggaatcc aggagcccag aggtggccgg gtaagaggcc tggtggtccc
                                                                         360
                                                                         420
47 ccactaaaag cctgcagtgt tcatgatcca actctcccta cagctccatg tcgctggatt
                                                                         480
49 ctcagectet gtgeettetg tetecacate tetetagaca gateteteac tgtetetagt
51 taggagteae tgtetetagt taggggtete tetgtetete tgaatetata tetecatgte
                                                                         540
53 taacteteag actgtetetg aggatatete teaageacte tgteteteeg getetgatte
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55 tetgtgtgte tteceteeat gettgtttgt gggtggetag acaccatete tececattea
                                                                         660
57 cagatggcta gatgctttct ctaaactttc ctttctacct agttctctct ctctcttt
                                                                         720
59 teceatetet etetetett ttetetetea gtetetaaat etgtetetet aggttetggg
                                                                         780
                                                                         840
61 tecatggatg ggagaggggg tagatggtet aggetettge etacetaata aegteecaga
                                                                         900
63 gggaagaaag ggagggacaa agagagggat ggagagactt gggctgaaga tccccagaca
                                                                         960
65 cggctaagtc tcagtcctca tccccaggtg ctgacgtgat ggccacagca ggaaatccct
67 ggggctggtt cetggggtac ctcatecttg gtgtegeagg tatetgagta tgegtgtgtg
                                                                        1020
69 tgtctgtccg tgcttggggg cacagtgttt gttaatgttc aggtgtgact cagtgtcctc
                                                                        1080
71 ttgcttgtga ctgcaaaget geetgtgaga eggtacegtg ttateegtee geeatggetg
                                                                        1140
73 tgcccctgca actccttgta tcgtggtaaa tttgtgtgtg gcagtgtgcc tgggtgtgtg
                                                                        1200
                                                                        1260
75 gttgtacctg tgagactctg acagtttgtg cctctgaata tctggtggag tgacaacagt
77 gtaatgatga tatggggaca ggggaagccg agggtgcagg agattgtgct tcctggggcg
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79	tgatccattg	ctgggaatct	gtgcctgctt	cctgggtctt	cagtcctgag	atccccctct	1380
81	cccatcccca	aggaactcac	ctcacaggac	tataaaacgg	tgttttggtg	tgcatgggct	1440
83	tgtggcttgg	tgtgactgtg	ggcaaggctg	ggagaggata	ggagtgactc	ggcgcaggac	1500
85	cgactctttg	agcatcagtc	tgcgcagaca	agtgacccga	tccttgctcc	cagcaacaac	1560
87	tccaccccct	gagctttaat	tcaccccgaa	ggacccgatc	ctaccgctat	gagcctagac	1620
89	tcctctgttg	aacccctcct	gaccgtggct.	ttgcaccgcg	atggcaccag	tctcacctcc	1680
		ccagagccct			•		1740
		taacccagct					1800
		tgatcccgcc					1860
		cctgaatccc					1920
		atcctggacc					1980
						teggeeeec	2040
						cctccccaa	2100
						aggagccctg	2160
						gaaccctggg	2220
						gtctctggta	2280
						tggcaggcgg	2340
						cagtgggtgc	2400
						gcagggcctg	2460
						aaacagcggt	2520
						ggaagggctg	2580
						ctcagcccag	2640
						aatgaggatt	2700
						tggggttagg	2760
						ttgccacaaa	2820
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						acggcaccca	2940
						cgaatccgtg	3000
						cgcggggaac	3060
						tgtgtgtctg	3120
						tgcgtcccag	3180
						gaggtctgca	3240
						gggcaagacc	3300
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						gaaacacaca	3540
						acctccaccc	3600
						tagcctactg	3660
157	ttgacgggga	gccttaccaa	taacataaat	agtcgattta	tgcatacgtt	ttatgcattc	3720
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161	tttttaaatt	gttgcaactc	tcctaaaatt	tttctgatgt	gtttattgaa	aaaatccaag	3840
163	tataagtgga	cttgtgcagt	tcaaaccagg	gttgttcaag	ggtcaactgt	gtacccagag	3900
165	ggaaacagtg	acacagatto	atagaggtga	aacacgaaga	gaaacaggaa	aaatcaagac	3960
167	tctacaaaga	ggctgggcag	ggtggctcat	gcctgtaato	ccagcacttt	gggaggcgag	4020
169	gcaggcagat	cacttgaggt	aaggagttca	agaccagcct	ggccaaaatg	gtgaaatcct	4080
171	gtctgtacta	aaaatacaaa	agttagctgg	atatggtggc	: aggcgcctgt	aatcccagct	4140
173	acttgggagg	ctgaggcagg	agaattgctt	gaatatggga	ı ggcagaggtt	gaagtgagtt	4200
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Input Set : A:\MTS3USA.seq.txt

Output Set: N:\CRF4\05092005\1936271C.raw

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177 aaaaaaaaag actttacaaa gagatgcaga gacactgaga cagataaaca agccacaaag
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4380
181 agcattcagg acataggaca tcgggaagca ggattagatg aagtcaggga tctggaatgg
                                                                       4440
183 gacttccaac agatatgttg ctgggctatg ttgttattga tgatggttct gtctttgttt
                                                                       4500
185 ctcagtctca tttagttcct ttctgagccc atatccattt ccacctctct gtgttttgaa
187 ttctgactct ccctctcttc acaacagggt gactctgggg ggcccctgat ctgcaacggg
                                                                       4620
189 tacttgcagg gccttgtgtc tttcggaaaa gccccgtgtg gccaagttgg cgtgccaggt
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191 gtctacacca acctctgcaa attcactgag tggatagaga aaaccgtcca ggccagttaa
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194 <210> SEQ ID NO: 2
195 <211> LENGTH: 237
196 <212> TYPE: PRT
197 <213> ORGANISM: Homo sapiens
199 <400> SEQUENCE: 2
201 Ser Leu Val Ser Gly Ser Cys Ser Gln Ile Ile Asn Gly Glu Asp Cys
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205 Ser Pro His Ser Gln Pro Trp Gln Ala Ala Leu Val Met Glu Asn Glu
206
               20
                                   25
209 Leu Phe Cys Ser Gly Val Leu Val His Pro Gln Trp Val Leu Ser Ala
213 Ala His Cys Phe Gln Asn Ser Tyr Thr Ile Gly Leu Gly Leu His Ser
217 Leu Glu Ala Asp Gln Glu Pro Gly Ser Gln Met Val Glu Ala Ser Leu
218 65
221 Ser Val Arg His Pro Glu Tyr Asn Arg Pro Leu Leu Ala Asn Asp Leu
                   85
                                       90
225 Met Leu Ile Lys Leu Asp Glu Ser Val Ser Glu Ser Asp Thr Ile Arg
               100
                                   105
229 Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr Ala Gly Asn Ser Cys Leu
           115
                               120
233 Val Ser Gly Trp Gly Leu Leu Ala Asn Gly Glu Leu Thr Gly Arg Met
234
       130
                           135
237 Pro Thr Val Leu Gln Cys Val Asn Val Ser Val Val Ser Glu Glu Val
                       150
                                           155
241 Cys Ser Lys Leu Tyr Asp Pro Leu Tyr His Pro Ser Met Phe Cys Ala
                   165
                                       170
245 Gly Gly Gln Asp Gln Lys Asp Ser Cys Asn Gly Asp Ser Gly Gly
246
                                   185
249 Pro Leu Ile Cys Asn Gly Tyr Leu Gln Gly Leu Val Ser Phe Gly Lys
250
           195
                               200
                                                   205
253 Ala Pro Cys Gly Gln Val Gly Val Pro Gly Val Tyr Thr Asn Leu Cys
                           215
257 Lys Phe Thr Glu Trp Ile Glu Lys Thr Val Gln Ala Ser
258 225
                       230
                                           235
261 <210> SEQ ID NO: 3
262 <211> LENGTH: 254
263 <212> TYPE: PRT
264 <213> ORGANISM: Homo sapiens
266 <400> SEQUENCE: 3
268 Met Ala Thr Ala Gly Asn Pro Trp Gly Trp Phe Leu Gly Tyr Leu Ile
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Input Set : A:\MTS3USA.seq.txt

Output Set: N:\CRF4\05092005\1936271C.raw

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10
272 Leu Gly Val Ala Gly Ser Leu Val Ser Gly Ser Cys Ser Gln Ile Ile
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276 Asn Gly Glu Asp Cys Ser Pro His Ser Gln Pro Trp Gln Ala Ala Leu
280 Val Met Glu Asn Glu Leu Phe Cys Ser Gly Val Leu Val His Pro Gln
284 Trp Val Leu Ser Ala Ala His Cys Phe Gln Asn Ser Tyr Thr Ile Gly
288 Leu Gly Leu His Ser Leu Glu Ala Asp Gln Glu Pro Gly Ser Gln Met
                    85
292 Val Glu Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Pro Leu
                                    105
296 Leu Ala Asn Asp Leu Met Leu Ile Lys Leu Asp Glu Ser Val Ser Glu
            115
                                120
300 Ser Asp Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr Ala
        130
                            135
                                                 140
304 Gly Asn Ser Cys Leu Val Ser Gly Trp Gly Leu Leu Ala Asn Gly Arg
                        150
                                             155
308 Met Pro Thr Val Leu Gln Cys Val Asn Val Ser Val Val Ser Glu Glu
                    165
                                         170
312 Val Cys Ser Lys Leu Tyr Asp Pro Leu Tyr His Pro Ser Met Phe Cys
                                     185
316 Ala Gly Gly Gly Gln Asp Gln Lys Asp Ser Cys Asn Gly Asp Ser Gly
317
            195
                                200
                                                     205
320 Gly Pro Leu Ile Cys Asn Gly Tyr Leu Gln Gly Leu Val Ser Phe Gly
                            215
324 Lys Ala Pro Cys Gly Gln Val Gly Val Pro Gly Val Tyr Thr Asn Leu
                        230
                                             235
328 Cys Lys Phe Thr Glu Trp Ile Glu Lys Thr Val Gln Ala Ser
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333 <211> LENGTH: 278
334 <212> TYPE: DNA
335 <213> ORGANISM: Homo sapiens
337 <400> SEQUENCE: 4
338 tgacccgctg taccacccca gcatgttctg cgccggcgga gggcaagacc agaaggactc
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340 etgeaacggt gaetetgggg ggeeeetgat etgeaacggg taettgeagg geettgtgte
                                                                          120
342 tttcggaaaa gccccgtgtg gccaagttgg cgtgccaggt gcctacacca acctctgcaa
                                                                          180
344 attcactgag tggatagaga aaaccgtcca ggccagttaa ctctggggac tgggaaccca
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346 tgaaattgac ccccaaatac atcctgcgga aggaattc
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349 <210> SEQ ID NO: 5
350 <211> LENGTH: 20
351 <212> TYPE: DNA
352 <213> ORGANISM: Artificial
354 <220> FEATURE:
355 <223> OTHER INFORMATION: primer
357 <400> SEQUENCE: 5
358 tgacccgctg taccacccca
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Input Set : A:\MTS3USA.seq.txt

Output Set: N:\CRF4\05092005\I936271C.raw

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	<211> LENGTH: 20	
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	<220> FEATURE:	
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	<400> SEQUENCE: 6	
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	<210> SEQ ID NO: 7	
	<211> LENGTH: 21	
	<212> TYPE: DNA	
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	<220> FEATURE:	
379	<223> OTHER INFORMATION: primer	
381	<400> SEQUENCE: 7	
382	ggtgatctgc gccctggtcc t	21
385	<210> SEQ ID NO: 8	
386	<211> LENGTH: 21	
387	<212> TYPE: DNA	
388	<213> ORGANISM: Artificial	
390	<220> FEATURE:	
391	<223> OTHER INFORMATION: primer	
393	<400> SEQUENCE: 8	
394	aggtgtccgg tggaggtggc a	21
397	<210> SEQ ID NO: 9	
398	<211> LENGTH: 18	
399	<212> TYPE: DNA	
400	<213> ORGANISM: Artificial	
402	<220> FEATURE:	
403	<223> OTHER INFORMATION: primer	
405	<400> SEQUENCE: 9	
406	tgcgcaagtt caccctca	18
409	<210> SEQ ID NO: 10	
410	<211> LENGTH: 19	
411	<212> TYPE: DNA	
412	<213> ORGANISM: Artificial	
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415	<223> OTHER INFORMATION: primer	
417	<400> SEQUENCE: 10	
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421	<210> SEQ ID NO: 11	
422	<211> LENGTH: 20	
423	<212> TYPE: DNA	
	<213> ORGANISM: Artificial	
426	<220> FEATURE:	
	<223> OTHER INFORMATION: primer	
	<400> SEQUENCE: 11	
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	<210> SEQ ID NO: 12	

Input Set : A:\MTS3USA.seq.txt

Output Set: N:\CRF4\05092005\1936271C.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

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DATE: 05/09/2005 TIME: 17:05:00

PATENT APPLICATION: US/09/936,271C

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Output Set: N:\CRF4\05092005\1936271C.raw